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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.						
09/809,638	03/14/2001	Mary Faris	s G&C 129.35-US-01 5083							
36327	7590 10/04/2004		EXAM	EXAMINÉR						
	C/O MORRISON & FOI	HARRIS, ALANA M ART UNIT PAPER NUMBER								
	EY CENTRE DRIVE, SUIT D. CA 92130									
5. 2. 2. 2. 2	, , ,		1642							
		DATE MAILED: 10/04/2004								

Please find below and/or attached an Office communication concerning this application or proceeding.

		Applicatio	n No.	Applicant(s)										
		09/809,63		FARIS ET AL.										
	Office Action Summary	Examiner		Art Unit										
		Alana M. I	farris, Ph.D.	1642										
	The MAILING DATE of this communication appears on the cover sheet with the correspondence address													
Period fo	• •													
THE External	ORTENED STATUTORY PERIOD FOR MAILING DATE OF THIS COMMUNICA nations of time may be available under the provisions of SIX (6) MONTHS from the mailing date of this communical period for reply specified above is less than thirty (30) of period for reply is specified above, the maximum statuture to reply within the set or extended period for reply will reply received by the Office later than three months after the patent term adjustment. See 37 CFR 1.704(b).	ATION. 7 CFR 1.136(a). In no eve cation. lays, a reply within the statu ory period will apply and will. by statute. cause the appli	nt, however, may a reply be tirr tory minimum of thirty (30) day expire SIX (6) MONTHS from cation to become ABANDONE	nely filed s will be considered time the mailing date of this o D (35 U.S.C. § 133).	ly. ommunication.									
Status														
1)[汉]	Responsive to communication(s) filed	on <i>14 June 2004</i> .												
,	•	This action is no	on-final.											
,	Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under <i>Ex parte Quayle</i> , 1935 C.D. 11, 453 O.G. 213.													
Disposit	ion of Claims													
4)⊠ 5)□ 6)⊠ 7)□	4) ☐ Claim(s) 1,14 and 23 is/are pending in the application. 4a) Of the above claim(s) is/are withdrawn from consideration. 5) ☐ Claim(s) is/are allowed. 6) ☐ Claim(s) 1,14 and 23 is/are rejected. 7) ☐ Claim(s) is/are objected to. 8) ☐ Claim(s) are subject to restriction and/or election requirement.													
Applicat	ion Papers													
9)[The specification is objected to by the E	Examiner.			·									
10)[The drawing(s) filed on is/are: a													
	Applicant may not request that any objection													
11)	Replacement drawing sheet(s) including the The oath or declaration is objected to be													
Priority (ınder 35 U.S.C. § 119													
 12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f). a) All b) Some * c) None of: 1. Certified copies of the priority documents have been received. 2. Certified copies of the priority documents have been received in Application No. 3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)). * See the attached detailed Office action for a list of the certified copies not received. 														
Attachmer	nt(s)													
1) 🛛 Notic	ce of References Cited (PTO-892)		4) Interview Summary											
3) 🔀 Infor	ce of Draftsperson's Patent Drawing Review (PTC mation Disclosure Statement(s) (PTC-1449 or PT er No(s)/Mail Date		Paper No(s)/Mail Do 5) Notice of Informal F 6) Other:		O-152)									

Art Unit: 1642

DETAILED ACTION

- 1. The finality of the action mailed May 19, 2004 has been withdrawn and PROSECUTION IS HEREBY REOPENED. A new action is set forth below.
- 2. Claims 1, 14 and 23 are pending.

Claim 7 has been cancelled.

Claims 14 and 23 have been amended.

Claims 1, 14 and 23 are examined on the merits.

Withdrawn Rejections

Claim Rejections - 35 USC § 112

- 3. The rejection of claims 14 and 23 are rejected under 35 U.S.C. 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention is withdrawn. Claim 7 has been cancelled.
- 4. The rejection of claims 14 and 23 under 35 U.S.C. 112, first paragraph, because the specification, does not reasonably provide enablement commensurate with the scope of the claimed invention is withdrawn. Claim 7 has been cancelled.

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New Grounds of Rejection

Claim Rejections - 35 USC § 102

5. The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

(e) the invention was described in a patent granted on an application for patent by another filed in the United States before the invention thereof by the applicant for patent, or on an international application by another who has fulfilled the requirements of paragraphs (1), (2), and (4) of section 371(c) of this title before the invention thereof by the applicant for patent.

The changes made to 35 U.S.C. 102(e) by the American Inventors Protection Act of 1999 (AIPA) and the Intellectual Property and High Technology Technical Amendments Act of 2002 do not apply when the reference is a U.S. patent resulting directly or indirectly from an international application filed before November 29, 2000. Therefore, the prior art date of the reference is determined under 35 U.S.C. 102(e) prior to the amendment by the AIPA (pre-AIPA 35 U.S.C. 102(e)).

6. Claims 2, 14 and 23 are rejected under 35 U.S.C. 102(e) as being anticipated by WO200270539 A2 (filed March 5, 2002/ IDS reference 2, April 15, 2001).
WO200270539 discloses sequence 1397, an amino acid sequence that is the same as Applicants' SEQ ID NO: 2, see the three attached database sheets.
The isolated 125P5C8 protein (SEQ ID NO: 2) encoded by a 125P5C8 nucleotide, where T can be U comprises a polynucleotide of at least 10 bases of Figure 2 (SEQ ID NO: 1) that comprises the base at position 2065. Absent evidence to the contrary the disclosed polynucleotide that encodes a 125P5C8

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protein identified as SEQ ID NO: 2 is contained in the plasmid designated Escherichia coli DH5A 125P5C8PRO deposited with American Type Culture Collection as Accession No. PTA-3137.

Double Patenting

7. The nonstatutory double patenting rejection is based on a judicially created doctrine grounded in public policy (a policy reflected in the statute) so as to prevent the unjustified or improper timewise extension of the "right to exclude" granted by a patent and to prevent possible harassment by multiple assignees. See *In re Goodman*, 11 F.3d 1046, 29 USPQ2d 2010 (Fed. Cir. 1993); *In re Longi*, 759 F.2d 887, 225 USPQ 645 (Fed. Cir. 1985); *In re Van Ornum*, 686 F.2d 937, 214 USPQ 761 (CCPA 1982); *In re Vogel*, 422 F.2d 438, 164 USPQ 619 (CCPA 1970);and, *In re Thorington*, 418 F.2d 528, 163 USPQ 644 (CCPA 1969).

A timely filed terminal disclaimer in compliance with 37 CFR 1.321(c) may be used to overcome an actual or provisional rejection based on a nonstatutory double patenting ground provided the conflicting application or patent is shown to be commonly owned with this application. See 37 CFR 1.130(b).

Effective January 1, 1994, a registered attorney or agent of record may sign a terminal disclaimer. A terminal disclaimer signed by the assignee must fully comply with 37 CFR 3.73(b).

8. Claims 1, 14 and 23 are provisionally rejected under the judicially created doctrine of obviousness-type double patenting as being unpatentable over claims 1, 12-17, 19, 21, 26, 51 and 52 of copending Application No. 10/099,460 (filed March 13, 2002). Although the conflicting claims are not identical, they are not patentably distinct from each other because SEQ ID NO: 2 of the instant application is 100% identical to SEQ ID NO: 3 and SEQ ID NO: 13 of the copending application reading on claim 1 of the copending application, 10/099,460 and intrinsically possessing all the attributes of the sequence disclosed in '460. Furthermore, SEQ ID NO: 1 of the instant application is

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the same as copending application's '460 nucleic acid sequence identified as SEQ ID NO: 12.

This is a <u>provisional</u> obviousness-type double patenting rejection because the conflicting claims have not in fact been patented.

- 9. Claims 1, 14 and 23 directed to an invention not patentably distinct from claims 1, 12-17, 19, 21, 26, 51 and 52 of commonly assigned co-pending application 10/099,460 (filed March 13, 2002). Specifically, both sets of claims read on identical nucleotide sequences and the encoded polypeptide sequence.
- 10. The U.S. Patent and Trademark Office normally will not institute an interference between applications or a patent and an application of common ownership (see MPEP § 2302). Commonly assigned 10/099,460 (filed March 13, 2002), discussed above, would form the basis for a rejection of the noted claims under 35 U.S.C. 103(a) if the commonly assigned case qualifies as prior art under 35 U.S.C. 102(e), (f) or (g) and the conflicting inventions were not commonly owned at the time the invention in this application was made. In order for the examiner to resolve this issue, the assignee can, under 35 U.S.C. 103(c) and 37 CFR 1.78(c), either show that the conflicting inventions were commonly owned at the time the invention in this application was made, or name the prior inventor of the conflicting subject matter.

A showing that the inventions were commonly owned at the time the invention in this application was made will preclude a rejection under 35 U.S.C. 103(a) based upon

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the commonly assigned case as a reference under 35 U.S.C. 102(f) or (g); or 35 U.S.C.

102(e) for applications filed on or after November 29, 1999.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACTTCACTAAACCCAATCTGGAGTTATCAGATGTCCAACAAGTGATGATACTGACATTAAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTCAGAATTTGGGGATTCATTTTAGGACAGATTGTTGTTGTTGTTGTTGTCTACGCATATGGTAT 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTCTACCATGACCTGGGACCGATGATCTATTACTTTCCTTTGCAAACACTAGAACTCACT 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCCATAGCCACACTTGATCGTATTGGCACAGATGGTGACTGCAGTAAACCTGAAGAAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTCAGAATTTGGGGATTCATTTTAGGACAGATTGTTCTTGTTCTACGCATATGGTAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TCTTCCTCACTGATAGTGCAAGCTGTGACTTGGTGGTCAGGAAGTCATTTGCAAAGGTAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TCTTCCTCACTGATAGTGCAAGCTGTGACTTGGTGGTGGGGAAGTCATTTGCAAAGGTAC 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGGCTTGAAGGTTTTAGTATAGCATTTCTTTCTCCCAATATTCCTAACAATTACTCCTTTC 362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGGCTTGAAGGTTTTAGTATAGCATTTCTTTCTCCAATATTCCTAACAATTACTCCTTTC 180
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1981 GACCACAGAGAAGTITCTGAGAAAATTCATTTAATCCCAGATTTGGATCCTACAAAA	301 AAGGATATCGACAGCACTGATGATGACAGATGGTGTGAATACATTATGTI	1621 CTGGTGGATTTTGTCGTGACACACTTTGGGAACACGAGAACATGACCTCGACAGACA	441 ATGTGGCTAGGGAAAAGTTGGGTTTCTATACAGACTTTGGTCCAAGCACAGGTP [[]]		263 GGGCTGAATATGCTATTTGGTCCTAAGAAAAACCTTGATTTGCTFCTTCAAACI 141 AGTTCTAAAGTGCTTTTCAGAAAGAGTGAAAAAATACATGAAACTTTTTCTGTGC 141 AGTTCTAAAGTGCTTTTCAGAAAGAGTGAAAAATACATGAAACTTTTTCTGTGC 323 AGTTCTAAAGTGCTTTTCAGGAAGAGTGAAAAATACATGAAAACTTTTTCTGTGC 324 AGTTCTAAAGTGCTTTTCAGGAAAGAGTGAAAAAATACATGAAAACTTTTTCTGTGC 325 AGTTCTAAAGTGCTTTTCAGGAATTAGGACTACGGCATAAAGCCTATGAGAGAAAA 386 GTTGGTGTGGGGATTGGTGTAGGGACTACGGCATAAAGCCTATGAGAGAAAA 387 GTTGGTGTGGGGATTGGTGGGGATTAGGACTACGGCATAAAGCCTATGAGAGAAAA
CCTACAAGAA 2040 CCTACAAGAA 2222 	TCCAGGGCTG 1860	CAGGAAACTG 1680	Hold Hold	rGAC 132 rGAT 138	AAAAAC 132 CTGCTT 120 CTGCTT 138 CTGGGC 126 [

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ABU43487	ADB65311	ADE40118	AAG59436	AAG58055	AAG59184	AAG59437	AAG58056	ABP58050	AAW94322	AAW55786	AAY38788	ABB90887	ABU32096	ABU24105	ABJ25677	ABU31622	AAE26672	ADA33802	ABM70704
Abu43487	Adb65311	Ade40118	Aag59436	Aag58055	Aag59184	Aag59437	Aag58056	Abp58050	Aaw94322	Aaw55786	Aay38788	Abb90887	Abu32096	Abu24105	Abj25677	Abu31622	. Aae26672	Ada33802	Abm70704
Protein e	Human pro	Human NOV	Arabidops	Arabidops	Arabidops	Arabidops	Arabidops	Human aqu	Human agu	Human aqu	Neisseria	Herbicida	Protein e	Protein e	Aspergill	Protein e	Human 599	Acinetoba	Photorhab

ALIGNMENTS

ABP69350 standard; protein; 699 Š

ABP69350;

20-JAN-2003 (first entry)

Human polypeptide SEQ ID NO 1397.

RESULT 1
ARBOJICO 1
AR Human; genome mapping; gene therapy; food supplement; virus; fungus; cell-proliferative disorder; neurodegenerative disease; bacterial; Parkinson's disease; Alzheimer's disease; autoimmune disease; multiple sclerosis; diabetes; genetic disorder; wound; burn; infection; arthritis; cytostatic; immunomodulator; nootropic; neuroprotective; antiparkinsonian; antidiabetic; immunosuppressive; dermatological; haemostatic; vulnerary; fungicide; antibacterial; virucide; protozoacide; antiarthritic.

Homo sapiens.

WO200270539-A2

12-SEP-2002

05-MAR-2002; 2002WO-US005095.

05-MAR-2001; 2001US-00799451.

(HYSE-) HYSEQ INC.

Tang YT, Zhou P, Goodrich RW, Asundi V, Zhang J, Xue AJ, Yang Y, Ma Y, Yamazaki V, Chen R, Wang Z, Wehrman T, Wang J, Wang D, Drmanac RT; Zhao QA, , Ghosh M; Ren F;

WPI; 2002-759812/82. N-PSDB; ABZ11567.

New polynucleotides comprising sequences assembled from expressed sequence tags (ESTs), useful for treating cell-proliferative, neurodegenerative, autoimmune, genetic, myeloid or lymphoid, or pineurodegenerative. or coagulation disorders. or platelet

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Claim 9; SEQ ID NO 1397; 1012pp + Sequence Listing; English.

The invention relates to an isolated polynucleotide (I) comprising a nucleotide sequence selected from any of 948 sequences (ABZ11119-ABZ12066) or their mature protein coding portion, active domain coding

Art Unit: 1642

Application/Control Number: 09/809,638

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	601 KDIDSTDHDRWCEYIMYRGLIRLGYARISHAELSDSEIQMAKFRIPDDPTNYRDNQKVVI 6	IDSTDHDRWCEYIMYRGLIRLGYARISHAELSDSEIQMAKFRIEDDFTNYRDNQKVVI	41 LVDEVVTHFGNHEDDLDRXLQALAVSKLLKSSSNQVIELGYITSAPGSRDYLQLTEHGNV		1	fgpstryhtwgimalsrypivksbhhllpspbgblapaitltvnisgk. 5	XVAPTKEVSAAIMPEREGYDNEGMSSLERSAHLLNETGADFITILESDASKEYMGNNDLT 4	PFRFGYDNEGWSSLERSAHLLNETGADFITILESDASKPYMGNNDLT 4	361 GLWMLFGFKONLDLLLQTKNSSKVLFRKSEKYMGCFLMLLVGVGLLGLGLRKAYERKLG 4	LLIOTKNISKVIJERKSEKYMKLEUMILVGVGLIGIGIRHKAVERKIG 4	301 TLGHLINSGTNPGKTWTIAMIFYLLEIFFCAWCTAFKFVPGGVYARERSDVLLGTWHLII 3	-1-1	241 VILCIASCIMIPSCIMPROTGIMMVTGTASAAGILVIHTWAAAVSGVVAIFTASWKPQ 3	SGLMLPSCLWFRGTGLIWWVTGTASAAGLLYLHTWAAAVSGCVFAIFTASMWPQ	181 KTGEVATGWASRPNWILAGAAFGSIVFITHWVFGEVSIVSRWAVSGHPHPGPDPNPFGGA 24	GEVATGMASRAMULAGAAFGSLVETTHWVFGEVSLVSKWAVSGHPHPGPDFNPFG	121 LRIWGFILGQIVLVVLRIWYTSLNPIWSYQWSNKVILTUSAIATLDRIGTDGDCSKPERK 18	IMGFILGQIVLVVLRIWYTSINPIWSYQMSNKVILTISAIATLDRIGTDGDCSKPEEK 1	61 WILVNEXWALTULRITTIGSTASPOAPNAKLELMYLALGVSSSLIVQAVYMYSGSKLQRY 1	61 WKLVNKKWMLTLLRIITIGSIASEQADNAKLRLMVIALGVSSSLIVQAVTWWSGSHLQRY 1	1 MTSLWREILLESLLGCVSWSLYHDLGPMIYYFPLQTLELTGLEGFSLAFLSPIFLTITPF 6	1 MTSLWREILLESULGCVSWSLYHIHIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII	Query Match 100.0%; Score 3720; DB 5; Length 699; Best Local Similarity 100.0%; Pred. No. 0; Matches 699; Conservative 0; Mismatches 0; Indels 0; Gaps	Sequence 699 AA;	protein or complementary sequences. The polynucleotides are useful for identifying expressed genes or for physical mapping of human genome. The encoded polypeptides (ABP68902-ABP69849) are useful as molecular weight markers, as a food supplement, for generating antibodies, in medical imaging, screening and diagnostic assays and for treating celliproliferative disorders (cancer), neurodegenerative diseases (Parkinson's or Alzheimer's disease), autoimmune diseases (multiple sclerosis, diabetes, lupus) genetic disorders, wound, burns, incision, ulcers, platelet or coagulation disorders, wound, burns, incision, ulcers, liver or lung fibrosis, infections (Dacterial, viral, fungal, parasitic), arthritis, etc. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from MIPO at ftp.wipo.int/pub/published_pct_sequences
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Art Unit: 1642

7. Any inquiry concerning this communication or earlier communications from the examiner should be directed to Alana M. Harris, Ph.D. whose telephone number is (571)272-0831. The examiner works a flexible schedule, however she can normally be reached between the hours of 6:30 am to 5:30 pm.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Jeffrey Siew can be reached on (571) 272-0787. The fax phone number for the organization where this application or proceeding is assigned is 703-872-9306.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

Alana M. Harris, Ph.D. 28 September 2004

ALANA M. HARRIS, PH.D. PRIMARY EXAMINER